

SEQUENCE LISTING

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<120> NOVEL G PROTEIN-COUPLED RECEPTOR
UP-REGULATED IN PROSTATE CANCER AND USES THEREOF

<130> 129.24USU1

<150> 60/157,902

<151> 1999-10-05

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Phe Ile Leu Ile Gly Leu Pro Gly Leu Glu Glu Ala Gln Phe Trp Leu	
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Ala Phe Pro Leu Cys Ser Leu Tyr Leu Ile Ala Val Leu Gly Asn Leu	
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Thr Ile Ile Tyr Ile Val Arg Thr Glu His Ser Leu His Glu Pro Met	
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tat ata ttt ctt tgc atg ctt tca ggc att gac atc ctc atc tcc acc	363
Tyr Ile Phe Leu Cys Met Leu Ser Gly Ile Asp Ile Leu Ile Ser Thr	
65 70 75	
tca tcc atg ccc aaa atg ctg gcc atc ttc tgg ttc aat tcc act acc	411
Ser Ser Met Pro Lys Met Leu Ala Ile Phe Trp Phe Asn Ser Thr Thr	
80 85 90	
atc cag ttt gat gct tgt ctg cta cag att ttt gcc atc cac tcc tta	459
Ile Gln Phe Asp Ala Cys Leu Leu Gln Ile Phe Ala Ile His Ser Leu	
95 100 105	

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Ser Gly Met Glu Ser Thr Val Leu Leu Ala Met Ala Phe Asp Arg Tyr	
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Val Ala Ile Cys His Pro Leu Arg His Ala Thr Val Leu Thr Leu Pro	
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Arg Val Thr Lys Ile Gly Val Ala Ala Val Val Arg Gly Ala Ala Leu	
145 150 155	
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Asn Ile Leu Ser His Ser Tyr Cys Leu His Gln Asp Val Met Lys Leu	
175 180 185	
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190 195 200 205	
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Ile Ser Ala Ile Gly Leu Asp Ser Leu Leu Ile Ser Phe Ser Tyr Leu	
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ctt att ctt aag act gtg ttg ggc ttg aca cgt gaa gcc cag gcc aag	843
Leu Ile Leu Lys Thr Val Leu Gly Leu Thr Arg Glu Ala Gln Ala Lys	
225 230 235	
gca ttt ggc act tgc gtc tct cat gtg tgt gct gtg ttc ata ttc tat	891
Ala Phe Gly Thr Cys Val Ser His Val Cys Ala Val Phe Ile Phe Tyr	
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gta cct ttc att gga ttg tcc atg gtg cat cgc ttt agc aag cgg cgt	939
Val Pro Phe Ile Gly Leu Ser Met Val His Arg Phe Ser Lys Arg Arg	
255 260 265	
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Asp Ser Pro Leu Pro Val Ile Leu Ala Asn Ile Tyr Leu Leu Val Pro	
270 275 280 285	
cct gtg ctc aac cca att gtc tat gga gtg aag aca aag gag att cga	1035
Pro Val Leu Asn Pro Ile Val Tyr Gly Val Lys Thr Lys Glu Ile Arg	
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cag cgc atc ctt cga ctt ttc cat gtg gcc aca cac gct tca gag ccc	1083
Gln Arg Ile Leu Arg Leu Phe His Val Ala Thr His Ala Ser Glu Pro	
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aacatttttg aagacagtat tcagaaaaaa aatttcctta ataaaaata caactcagat	1203
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tataattatt	aatactctga	ctaggttggtg	ggttgagggt	tattactttt	cattttacca	240
tgcagtccaa	atctaaactg	cttctactga	tggtttacag	cattctgaga	taagaatggt	300
acatctagag	aacatttgcc	aaaggcctaa	gcacagcaaa	ggaaaataaa	cacagaatat	360
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tct tat atc ttt atc ctt cag gca gtt cta caa ctc tcc tct cag gag 96
Ser Tyr Ile Phe Ile Leu Gln Ala Val Leu Gln Leu Ser Ser Gln Glu
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gcc cgc tac aaa gca ttt ggg aca tgt gtc tct cac ata ggt gcc atc 144
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tta gcc ttc tac aca cct tca gtc atc tct tca gtc atg cac cgt gtg 192
Leu Ala Phe Tyr Thr Pro Ser Val Ile Ser Ser Val Met His Arg Val
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Ala Arg Cys Ala Val Pro His Val His Ile Leu Leu Ala Asn Phe Tyr
65 70 75 80

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Leu Leu Phe Pro Pro Met Val Asn Pro Ile Ile Tyr Gly Val Lys Thr
85 90 95

aag cag atc cgt gac agt ctt ggg agt att cct gag aaa gga tgt gtg 336
Lys Gln Ile Arg Asp Ser Leu Gly Ser Ile Pro Glu Lys Gly Cys Val
100 105 110

aat aga gag tga gga ata agt gga aaa aga gtg ggg ccc agt gaa tgc 384
Asn Arg Glu * Gly Ile Ser Gly Lys Arg Val Gly Pro Ser Glu Cys
115 120 125

tgt agt ggg cca ggg ctg tgc tga gag tag atg ggt cct aga ctc cac 432
Cys Ser Gly Pro Gly Leu Cys * Glu * Met Gly Pro Arg Leu His
130 135 140

gtt tag ttc ttt tct tgt att atg aaa aga ata aat gat gtc ctg aag 480
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Ala Arg Tyr Lys Ala Phe Gly Thr Cys Val Ser His Ile Gly Ala Ile
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Leu Ala Phe Tyr Thr Pro Ser Val Ile Ser Ser Val Met His Arg Val
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Ala Arg Cys Ala Val Pro His Val His Ile Leu Leu Ala Asn Phe Tyr

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